## Replacement Sheet 1/2 Application No. 10/650,559

11 10	7 7 9	7 7 9	7 7 0	7 7 0	7 7	7 7 9
NO: NO: NO:	NO: NO:	NO: NO:	NO: NO:	NO: NO: NO:	NO: NO: NO:	NO: NO: NO:
UD UD UD	ID ID ID	di di di	ID ID ID			
SEQ $SEQ$	SEQ SEQ SEQ	SEQ SEQ SEQ	SEQ SEQ SEQ	$\begin{array}{c} SEQ \\ SEQ \\ SEQ \end{array}$	SEQ SEQ SEQ	SEQ SEQ SEQ
YANCTNTGYA YANCTNTGYA YANCTNTGYA	100 HVAIFDSDAG HVAIFDSDAG	150 AVTVIGNIGL AVTVIGNIGL AVTVTDNTGL	200 SDRHQQQANT SDRHQQQANT SDRHQQQANT	250 DYEDSASADK DYEDSASADK DYEDSASADK	300 AKYPNSIWIA AKYPNSIWIA AKYPNSIWIA	350 VLLADDSSKV VLLADDSSKV VLLADDSSKM
ADYCKYLGLP ADYCKYLGLP ADYCKYLGLP	MVVAGVTPYS MVVAGVTPYS MVVDGVTPYS	TAFRPKVFKN TAFRPKVFKN TAFRPKSFKN	IKVSESLAWL IKVSESLAWL IKVSESLAWL	PTKKVSYLVI PTKKVSYLVI PSKKVSYLVI	TYNATDYQYII LINNIDYQYII LINNIDYQXII	GVAGGLDKNI GVAGGLDKNI GVAGGLDKNI
SLGAQCWDGY SLGAQCWDGY SFGAQCWDGY	TMQAGDVAIF TMQAGDVAIF VMQAGDVAIF	VKIPYSTTYP VKIPYSTTYP VKIPYSATYP	CQQAGTTKTI CQQAGTTKTI CQQAGTTKTI	RVADLFLSNL READLFLSNL READLFLSNL	PVYYSYKPFT PVYYSYKPFT PIYYSYKPFT	GVRWWQFTSV GVRWWQFTSV GVRWWQFTSV
SNGNAYDIDG SNGNAYDIDG SNGNAYDIDG	GILNYFDEVE GILNYFDEVE GILNYFDEVE	QGGANGAYNL QGGANGAYNL QGGANGAYNL	AYQQADLTAT AYQQADLTAT AYQQADLTTT	RFGGDSNLAQ RFGGDSNLAQ RFGGDSALAQ	MDKIANAGYK MDKIANAGYK MDKIASAGYK	PLWEFFPSMD PLWEFFPSMD PLWEFFPSMD
MATYQEYKSR MATYQEYKSR MATYQEYKSR	51 RDIWEQRHKN KDIWEQRHKN RDIWEQRHEN	101 SGYGWFLGQN SGYGWFLGQN GGYGWFLGQN	151 NKGDYFIDVS NKGDYFIDVS NKGDYFIDVS	201 SDPIGYYHFG SDPIGYYHFG SDPIGYYHFG	251 EANTKAVIAF EANTNAVIAF QANTNAVIAF	301 GYPDYEVRTE GYPDYEVRTE GYPDYEVRTE
lysin lysin lysin	lysin Iysin Iysin	lysin lysin lysin	lysin lysin lysin	lysin lysin lysin	lysin lysin lysin	lysin lysin lysin
group C phage group A phage group B phage	group C phage group A phage group B phage	group C phage group A phage group B phage				

## Replacement Sheet 2/2 Application No. 10/650,559